



IFW16

## RAW SEQUENCE LISTING

DATE: 08/26/2004

PATENT APPLICATION: US/09/723,722B

TIME: 11:14:33

Input Set : A:\152706441USSEQ3.txt

Output Set: N:\CRF4\08262004\I723722B.raw

4 <110> APPLICANT: Anderson, John P.  
 5 Basi, Gurigbal  
 6 Doane, Minh Tam  
 7 Frigon, Normand  
 8 John, Varghese  
 9 Power, Michael  
 10 Sinha, Sukanto  
 11 Tatsuno, Gwen  
 12 Tung, Jay  
 13 Wang, Shuwen  
 14 McConlogue, Lisa

16 <120> TITLE OF INVENTION: Beta-Secretase Enzyme Compositions and  
 17 Methods

19 <130> FILE REFERENCE: 228-US-NEWC1

21 <140> CURRENT APPLICATION NUMBER: 09/723,722B

22 <141> CURRENT FILING DATE: 2000-11-28

24 <150> PRIOR APPLICATION NUMBER: US 09/501,708

25 <151> PRIOR FILING DATE: 2000-02-10

27 <150> PRIOR APPLICATION NUMBER: 60/119,571

28 <151> PRIOR FILING DATE: 1999-02-10

30 <150> PRIOR APPLICATION NUMBER: 60/139,172

31 <151> PRIOR FILING DATE: 1999-06-15

33 <160> NUMBER OF SEQ ID NOS: 104

35 <170> SOFTWARE: FastSEQ for Windows Version 4.0

37 <210> SEQ ID NO: 1

38 <211> LENGTH: 1503

39 <212> TYPE: DNA

40 <213> ORGANISM: Homo sapiens

42 <400> SEQUENCE: 1

43 atggcccaag cctgcctg gctcctgctg tggatgggag cgggagtgt gctgcccac 60  
 44 ggcacccagc acggcatcgg gctgcccctg cgcagcggcc tggggggcgc ccccttggg 120  
 45 ctgctggctgc cccgggagac cgacgaagag cccgaggagc ccggccggag gggcagcttt 180  
 46 gtggagatgg tggacaacct gaggggcaag tcggggcagg gctactacgt ggagatgacc 240  
 47 gtgggcagcc ccccgagac gctcaacatc ctggtggata caggcagcag taactttgca 300  
 48 gtgggtgctg cccccaccc cttcctgcat cgctactacc agaggcagct gtccagcaca 360  
 49 taccgggacc tccggaagg tgtgtatgtg ccctacaccc agggcaagt ggaaggggag 420  
 50 ctgggcaccg acctggttaag catccccat ggccccaaag tcaactgtgc tgccaacatt 480  
 51 gctgccatca ctgaatcaga caagtcttc atcaacggct ccaactggga aggcacctg 540  
 52 gggctggcct atgctgagat tgccaggcct gacgactccc tggagccttt ctttgactct 600  
 53 ctggttaaagc agaccacgt tcccaacctc ttctccctgc agctttgtgg tgetggttc 660  
 54 cccctcaacc agtctgaagt gctggcctct gtcggaggga gcatgatcat tggaggtatc 720  
 55 gaccactcgc tgtacacagg cagtctctgg tatacaccca tccggcggga gtggtattat 780  
 56 gaggtgatca ttgtgcgggt ggagatcaat ggacaggatc tgaaaatgga ctgcaaggag 840

(pg. 6)

**ENTERED**

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57 tacaactatg acaagagcat tgtggacagt ggcaccacca accttcgttt gcccagaaa      900
58 gtgtttgaag ctgcagtcaa atccatcaag gcagcctcct ccacggagaa gttccctgat      960
59 ggtttctggc taggagagca gctgggtgtgc tggcaagcag gcaccacccc ttggaacatt     1020
60 ttcccagtc tctcactcta cctaattgggt gaggttacca accagtcctt ccgcatcacc     1080
61 atccttcgc agcaatacct gcggccagtg gaagatgtgg ccacgtccca agacgactgt     1140
62 tacaagtttg ccatctcaca gtcattccac ggcactgtta tgggagctgt tatcatggag     1200
63 ggcttctacg ttgtctttga tcgggcccga aaacgaattg gctttgctgt cagcgcttgc     1260
64 catgtgcacg atgagttcag gacggcagcg gtggaaggcc cttttgtcac cttggacatg     1320
65 gaagactgtg gctacaacat tccacagaca gatgagtcaa ccctcatgac catagcctat     1380
66 gtcattggctg ccatctgcgc cctcttcatt ctgccactct gcctcatggt gtgtcagtgg     1440
67 cgctgcctcc gctgcctgcg ccagcagcat gatgactttg ctgatgacat ctccctgctg     1500
68 aag                                          1503

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70 &lt;210&gt; SEQ ID NO: 2

71 &lt;211&gt; LENGTH: 501

72 &lt;212&gt; TYPE: PRT

73 &lt;213&gt; ORGANISM: Homo sapiens

75 &lt;400&gt; SEQUENCE: 2

```

76 Met Ala Gln Ala Leu Pro Trp Leu Leu Leu Trp Met Gly Ala Gly Val
77 1          5          10          15
78 Leu Pro Ala His Gly Thr Gln His Gly Ile Arg Leu Pro Leu Arg Ser
79          20          25          30
80 Gly Leu Gly Gly Ala Pro Leu Gly Leu Arg Leu Pro Arg Glu Thr Asp
81          35          40          45
82 Glu Glu Pro Glu Glu Pro Gly Arg Arg Gly Ser Phe Val Glu Met Val
83          50          55          60
84 Asp Asn Leu Arg Gly Lys Ser Gly Gln Gly Tyr Tyr Val Glu Met Thr
85 65          70          75          80
86 Val Gly Ser Pro Pro Gln Thr Leu Asn Ile Leu Val Asp Thr Gly Ser
87          85          90          95
88 Ser Asn Phe Ala Val Gly Ala Ala Pro His Pro Phe Leu His Arg Tyr
89          100         105         110
90 Tyr Gln Arg Gln Leu Ser Ser Thr Tyr Arg Asp Leu Arg Lys Gly Val
91          115         120         125
92 Tyr Val Pro Tyr Thr Gln Gly Lys Trp Glu Gly Glu Leu Gly Thr Asp
93          130         135         140
94 Leu Val Ser Ile Pro His Gly Pro Asn Val Thr Val Arg Ala Asn Ile
95 145         150         155         160
96 Ala Ala Ile Thr Glu Ser Asp Lys Phe Phe Ile Asn Gly Ser Asn Trp
97          165         170         175
98 Glu Gly Ile Leu Gly Leu Ala Tyr Ala Glu Ile Ala Arg Pro Asp Asp
99          180         185         190
100 Ser Leu Glu Pro Phe Phe Asp Ser Leu Val Lys Gln Thr His Val Pro
101          195         200         205
102 Asn Leu Phe Ser Leu Gln Leu Cys Gly Ala Gly Phe Pro Leu Asn Gln
103          210         215         220
104 Ser Glu Val Leu Ala Ser Val Gly Gly Ser Met Ile Ile Gly Gly Ile
105 225         230         235         240
106 Asp His Ser Leu Tyr Thr Gly Ser Leu Trp Tyr Thr Pro Ile Arg Arg
107          245         250         255

```

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```

108 Glu Trp Tyr Tyr Glu Val Ile Ile Val Arg Val Glu Ile Asn Gly Gln
109           260           265           270
110 Asp Leu Lys Met Asp Cys Lys Glu Tyr Asn Tyr Asp Lys Ser Ile Val
111           275           280           285
112 Asp Ser Gly Thr Thr Asn Leu Arg Leu Pro Lys Lys Val Phe Glu Ala
113           290           295           300
114 Ala Val Lys Ser Ile Lys Ala Ala Ser Ser Thr Glu Lys Phe Pro Asp
115 305           310           315           320
116 Gly Phe Trp Leu Gly Glu Gln Leu Val Cys Trp Gln Ala Gly Thr Thr
117           325           330           335
118 Pro Trp Asn Ile Phe Pro Val Ile Ser Leu Tyr Leu Met Gly Glu Val
119           340           345           350
120 Thr Asn Gln Ser Phe Arg Ile Thr Ile Leu Pro Gln Gln Tyr Leu Arg
121           355           360           365
122 Pro Val Glu Asp Val Ala Thr Ser Gln Asp Asp Cys Tyr Lys Phe Ala
123           370           375           380
124 Ile Ser Gln Ser Ser Thr Gly Thr Val Met Gly Ala Val Ile Met Glu
125 385           390           395           400
126 Gly Phe Tyr Val Val Phe Asp Arg Ala Arg Lys Arg Ile Gly Phe Ala
127           405           410           415
128 Val Ser Ala Cys His Val His Asp Glu Phe Arg Thr Ala Ala Val Glu
129           420           425           430
130 Gly Pro Phe Val Thr Leu Asp Met Glu Asp Cys Gly Tyr Asn Ile Pro
131           435           440           445
132 Gln Thr Asp Glu Ser Thr Leu Met Thr Ile Ala Tyr Val Met Ala Ala
133           450           455           460
134 Ile Cys Ala Leu Phe Met Leu Pro Leu Cys Leu Met Val Cys Gln Trp
135 465           470           475           480
136 Arg Cys Leu Arg Cys Leu Arg Gln Gln His Asp Asp Phe Ala Asp Asp
137           485           490           495
138 Ile Ser Leu Leu Lys
139           500

```

141 &lt;210&gt; SEQ ID NO: 3

142 &lt;211&gt; LENGTH: 24

143 &lt;212&gt; TYPE: DNA

144 &lt;213&gt; ORGANISM: Homo sapiens

146 &lt;400&gt; SEQUENCE: 3

147 gagagacgar garccwgagg agcc

24

149 &lt;210&gt; SEQ ID NO: 4

150 &lt;211&gt; LENGTH: 24

151 &lt;212&gt; TYPE: DNA

152 &lt;213&gt; ORGANISM: Artificial Sequence

154 &lt;220&gt; FEATURE:

155 &lt;223&gt; OTHER INFORMATION: Degenerate oligonucleotide primer derived from SEQ

156 ID NO: 2

158 &lt;400&gt; SEQUENCE: 4

159 gagagacgar garccwgaag agcc

24

161 &lt;210&gt; SEQ ID NO: 5

162 &lt;211&gt; LENGTH: 24

## RAW SEQUENCE LISTING

DATE: 08/26/2004

PATENT APPLICATION: US/09/723,722B

TIME: 11:14:33

Input Set : A:\152706441USSEQ3.txt

Output Set: N:\CRF4\08262004\I723722B.raw

163 <212> TYPE: DNA  
164 <213> ORGANISM: Artificial Sequence  
166 <220> FEATURE:  
167 <223> OTHER INFORMATION: Degenerate oligonucleotide primer derived from SEQ  
168 ID NO: 2  
170 <400> SEQUENCE: 5  
171 gagagacgar garccwgaag aacc 24  
173 <210> SEQ ID NO: 6  
174 <211> LENGTH: 24  
175 <212> TYPE: DNA  
176 <213> ORGANISM: Artificial Sequence  
178 <220> FEATURE:  
179 <223> OTHER INFORMATION: Degenerate oligonucleotide primer derived from SEQ  
180 ID NO: 2  
182 <400> SEQUENCE: 6  
183 gagagacgar garccwgagg aacc 24  
185 <210> SEQ ID NO: 7  
186 <211> LENGTH: 23  
187 <212> TYPE: DNA  
188 <213> ORGANISM: Artificial Sequence  
190 <220> FEATURE:  
191 <223> OTHER INFORMATION: Degenerate oligonucleotide primer derived from SEQ  
192 ID NO: 2  
194 <400> SEQUENCE: 7  
195 agagacgarg arccsgagga gcc 23  
197 <210> SEQ ID NO: 8  
198 <211> LENGTH: 23  
199 <212> TYPE: DNA  
200 <213> ORGANISM: Artificial Sequence  
202 <220> FEATURE:  
203 <223> OTHER INFORMATION: Degenerate oligonucleotide primer derived from SEQ  
204 ID NO: 2  
206 <400> SEQUENCE: 8  
207 agagacgarg arccsgaaga gcc 23  
209 <210> SEQ ID NO: 9  
210 <211> LENGTH: 23  
211 <212> TYPE: DNA  
212 <213> ORGANISM: Artificial Sequence  
214 <220> FEATURE:  
215 <223> OTHER INFORMATION: Degenerate oligonucleotide primer derived from SEQ  
216 ID NO: 2  
218 <400> SEQUENCE: 9  
219 agagacgarg arccsgaaga acc 23  
221 <210> SEQ ID NO: 10  
222 <211> LENGTH: 23  
223 <212> TYPE: DNA  
224 <213> ORGANISM: Artificial Sequence  
226 <220> FEATURE:  
227 <223> OTHER INFORMATION: Degenerate oligonucleotide primer derived from SEQ

## RAW SEQUENCE LISTING

DATE: 08/26/2004

PATENT APPLICATION: US/09/723,722B

TIME: 11:14:33

Input Set : A:\152706441USSEQ3.txt

Output Set: N:\CRF4\08262004\I723722B.raw

228 ID NO: 2  
230 <400> SEQUENCE: 10  
231 agagacgarg arccsgagga acc 23  
233 <210> SEQ ID NO: 11  
234 <211> LENGTH: 23  
235 <212> TYPE: DNA  
236 <213> ORGANISM: Artificial Sequence  
238 <220> FEATURE:  
239 <223> OTHER INFORMATION: Degenerate oligonucleotide primer derived from SEQ  
240 ID NO: 2  
242 <400> SEQUENCE: 11  
243 cgtcacagrt trtcaaccat ctc 23  
245 <210> SEQ ID NO: 12  
246 <211> LENGTH: 23  
247 <212> TYPE: DNA  
248 <213> ORGANISM: Artificial Sequence  
250 <220> FEATURE:  
251 <223> OTHER INFORMATION: Degenerate oligonucleotide primer derived from SEQ  
252 ID NO: 2  
254 <400> SEQUENCE: 12  
255 cgtcacagrt trtctaccat ctc 23  
257 <210> SEQ ID NO: 13  
258 <211> LENGTH: 23  
259 <212> TYPE: DNA  
260 <213> ORGANISM: Artificial Sequence  
262 <220> FEATURE:  
263 <223> OTHER INFORMATION: Degenerate oligonucleotide primer derived from SEQ  
264 ID NO: 2  
266 <400> SEQUENCE: 13  
267 cgtcacagrt trtccaccat ctc 23  
269 <210> SEQ ID NO: 14  
270 <211> LENGTH: 23  
271 <212> TYPE: DNA  
272 <213> ORGANISM: Artificial Sequence  
274 <220> FEATURE:  
275 <223> OTHER INFORMATION: Degenerate oligonucleotide primer derived from SEQ  
276 ID NO: 2  
278 <400> SEQUENCE: 14  
279 cgtcacagrt trtcgaccat ctc 23  
281 <210> SEQ ID NO: 15  
282 <211> LENGTH: 23  
283 <212> TYPE: DNA  
284 <213> ORGANISM: Artificial Sequence  
286 <220> FEATURE:  
287 <223> OTHER INFORMATION: Degenerate oligonucleotide primer derived from SEQ  
288 ID NO: 2  
290 <400> SEQUENCE: 15  
291 cgtcacagrt trtcaaccat ttc 23  
293 <210> SEQ ID NO: 16

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/723,722B

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Input Set : A:\152706441USSEQ3.txt  
Output Set: N:\CRF4\08262004\I723722B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:22; N Pos. 12  
Seq#:23; N Pos. 12  
Seq#:24; N Pos. 12  
Seq#:25; N Pos. 12  
Seq#:26; N Pos. 7  
Seq#:27; N Pos. 7  
Seq#:28; N Pos. 3,12  
Seq#:29; N Pos. 3,12  
Seq#:34; N Pos. 16  
Seq#:35; N Pos. 16  
Seq#:36; N Pos. 16  
Seq#:37; N Pos. 16  
Seq#:48; N Pos. 6164,6238,6254,6255,6256,6257,6258,6259,6260,6261,6262,6263  
Seq#:48; N Pos. 6264,6265,6266,6267,6268,6269,6270,6271,6272  
Seq#:61; Xaa Pos. 4  
Seq#:72; Xaa Pos. 10  
Seq#:73; Xaa Pos. 5  
Seq#:76; N Pos. 6,18,27,30,33,36,39,42,48,57  
Seq#:78; Xaa Pos. 3  
Seq#:81; Xaa Pos. 4

## VERIFICATION SUMMARY

DATE: 08/26/2004

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Input Set : A:\152706441USSEQ3.txt

Output Set : N:\CRF4\08262004\I723722B.raw

L:379 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0  
L:395 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0  
L:411 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0  
L:427 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0  
L:443 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0  
L:459 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0  
L:475 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0  
L:491 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0  
L:551 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0  
L:567 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0  
L:583 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:0  
L:599 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:0  
L:960 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48 after pos.:6120  
L:961 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48 after pos.:6180  
L:962 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48 after pos.:6240  
L:1475 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:0  
L:1967 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:72 after pos.:0  
L:1984 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:73 after pos.:0  
L:2112 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:76 after pos.:0  
L:2140 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78 after pos.:0  
L:2183 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81 after pos.:0